

Amendments to the Specification:

Please replace the paragraph beginning at page 6, line 15, with the following rewritten paragraph:

-- Particularly conserved regions and amino acid residues common to nectin polypeptides were identified by aligning nectin polypeptide sequences with each other and additional closely-related members of the nectin-Ig superfamily of proteins. The amino acid sequence of nectin-3 α and nectin-4 (SEQ ID Nos: 6 and 24) were compared with the amino acid sequences of other nectin and Ig family members (SEQ ID NO:20, 22, and 25), using a multiple sequence alignment program. The alignment of these sequences is shown in Table 2, and includes consensus residues (capitalized), which are identical among at least a majority [(three)] of the [five] amino acid sequences in the alignment. [In addition, lower case residues are shown on a separate line of Table 2 and represent residues that are not consensus residues, but are identical between human nectin-3 α and human nectin-4 (SEQ ID Nos: 6 and 24).] - -

Please replace Table 2 beginning at page 7, line 7, with the following rewritten Table:

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Table 2
Conserved Nectin Amino Acids

(Hs=Homo sapiens)
(Mus=Murine)

HUNECTIN2 (SEQ ID NO:22) HUCD155 (SEQ ID NO:25) HUNECTIN1 (SEQ ID NO:20) HUNECTIN3 (SEQ ID NO:6) HUNECTIN4 (SEQ ID NO:24) consensus	~~~~~ MARAAALLPS RSPPTPLLWP LLLLLLL... ~~~~~ MARAMAAAWP LLLVALLVLS ~~~~~MARMG LAGAAGRWWG L...ALGLTA MARTLRPSPL CPGGGKAQLS SASLLGAGLL LQPPTPPPLL LLLFPLLLFS ~~~~~MPLSLG AEMWGPEAWL LLLLLLASFT P W LLL LL
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	51 100 ..ETGAQDVR VQVLPEVRGQ LGGTVELPCH L.LPPVPGLY ISLVTWQRPD WPPPGTGDVV VQAPTQVPGF LGDSVTLPCY LQVPNMEVTH VSQLTWAR.. FFLPGVHSQV VQVNDSMYGF IGTDVVLHCS FANP.LPSVK ITQVTWQK.S RLCGALAGP. IIVEPHVTAV WGKNVSLKCL I..EV..NET ITQISWEKIH GRCP..AGE. LETSDVVTVV LGQDAKLPCF YRGDS..GEQ VGQVAWARVD -ePG-ag-VQV-VtGv-LG-V-LPC-P-e-I-QV-W-R PG VOV V G LG V LPC P I QVTW R
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	101 150 APANHQNVAA FHPKMGPSFP SPKPGSERLS FVSAKQSTGQ DTEAELQDAT .HGESGSMAV FHQTQGPSYS E....SKRLE FVAARLG...AELRNAS TNGSKQNVAI YNPSMGVSV. .LAPYRERVE FL.....RPSFTDGT .GKSSQTVAV HHPQYGFSVQ ..GEYQGRVL FKNYSLN...DAT AGEGAQELAL LHSKYGLHVS ..PAYEGRVE QPPPPRNPL.DGS -g-Q-A-H-yG-SV-Y-grVE-F-n-DAT Q VA HP G SV Y RVE E DAT

HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	151 LALHGLTVED EGNYTCEFAT FPKGSVRGMT WLRVIAKPKN QAEAQKVTF. LRMFGLRVED EGNYTCLFVT FPQGSRSVDI WLRVLAKPQN TAEVQKVQL. IRLSRLELED EGVYICEFAT FPTGNRESQL NLTVMAPPTN WIEGTQAVLR ITLHNIGFSD SGKYICKAVT FPLGNAQSST TVTVLVEPTV SLIKGPDSL VLLRNAVQAD EGEYECRVST FPAGSFQARL RLRVLVPPLP SLNPGP.ALE L nL ED EG Y C F T FP GS q LRVLAKP N s E L <u>L L ED EG Y C F T FP GS LRVLAKP N E L</u>	200
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	201SQDPTT VALCISKEGR PPARISWLSS LDWEAKETQV SGTLAGTVTVTGEPVP MARCVSTGGR PPAQITWHSD LGGMPNTSQV PGFLSGTVTV AKKGQDDKVL VATCTSANGK PPSVSVWETR LKGEARVPGD SGTPMAPVTV DGGNE...TV AAICIAATGK PVAHIDWEGD LGEM..ESTT TSFPNETATI EGQGL...TL AASC.TAEGS PAPSVTWDE VKGT..TSSR SFKHSRSAAV g T aA C Sa G PPA I W L G S SG TVTV <u>T A C SA G PPA I W L G S SG TVTV</u>	250
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	251 TSRFTLVPSG RADGVTVTCK VEH..ESFEE PALIPVTLVS RYPPEVSISG TSLWILVPSS QVDGKNVTCK VEH..ESFEK PQLLTVNLTV YYPPEVSISG ISRYRLVPSR EAHQQSLACI VNYHMDRFKESLTLNV QYEPEVTIEG ISQYKLFPTR FARGRRITCV VKHP..ALEK DIRYSFILDI QYAPEVSVTG TSEFHLVPSR SMNGQPLTCV VSHP..GLLQ DQRITHILHV SFLAEASVRG TS LVPSR A G TC V Hp FE d r iL V Y PEVSI G <u>TS LVPSR A G TC V H FE L V Y PEVSI G</u>	300
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	301 Y.DDN.WYLG RTDATLSCDV RSNPEPTGYD WSTTSGTFPT SAVAQGSQVLV Y.DNN.WYLG QNEATLTCD A RSNPEPTGYN WSTTMGPLPP FAVAQGAQLL F.DGN.WYLG RMDVKLTCKA DANPPATEYH WTTLNGSLPK GVEAQNRTLF Y.DGN.WFVG RKGVNLCNA DANPPPFKSV WSRLDGQWPD GLLASDNTLH LEDQNLWHIG REGAMLKCLS EGQPPPSYN. WTRLDGPLPS GVRVDGDTLG Y D N WYLG R gA LkC A NPPPTY WSTLdG LP G AOG TL <u>Y D N WYLG R A L C A NPPPT Y WSTL G LP G AOG TL</u>	350
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	351 IH.AVDSLFN TTFVCTVTNA VGMGRAEQVI FVRETP.... IR.PVDKPIN TTLICNVNTA LGARQAELTV QVKEGP.... FKGPINYSLA GTYICEATNP IGTRSGQVEV NITEFPYTPS FVHPLTFNYS GVIYICKVTNS LGQRSDQKVI YISDPPTTTT LQPTIQWHPS F.PPLTTEHS GIYVCHVSNE FSSRDSQVTV DVLDPQEDSG KQ..... F Plt s G YIC VTN G R Q V BpP q <u>F P GTYIC VTN G R Q V V E P</u>	400
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	401RAS P...RDV..G PLVWGAVGGT LLVLLLLAGGPSE H...SGISRN AIIFLVLG.. ILVFLILLGIPPE HGRRAGPVPT AIIGGVAGSI LLVLIVVGGI TADIEDLATE PKKLPFPLST LATIKDDTIA TIIASVVGGA LFIVLVSVLAVDLV..SAS VVVVGVI AAL LFCLLVVVV d II GV G LLVLLV vG <u>II GV G LLVLLV G</u>	450
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	451 SLAFILLRVR RR....RKS .PGGAGGGAS GDGGFYDPKA QVLGNQDPVF GIYFYWSKCS REVLWHCHLC .PSSEHHQSC RN~~~~~ VVALRRRRHT FKGDYSTKKH .VYGNGYSKA GIPQHPPMA QNLQYPDDSD GIFCYRRRT FRGDYFAKNY IPPSDMQES QIDVLQOQDEL D..SYP.DSV LMSRYHRR.. .KAQOMTQKY EEELTLTREN SIRRLHSHHT DPRSQPEESV y RR y P e I lH d Ls PD Sv <u>Y RR P I P S</u>	500

HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	501					550
	WTPVVPGPME	P.DGKDEEEE	EEEEKA EKGL	MLPPPPALED	DMESQLDGSL	
	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~	
	.DEKKAGPLG	G.SSYEEEE	EEEGGGGER	KVG GPHPKYD	EDAKR PYFTV	
	.KKENKNP..	.VNNLIRKDY	LEEPEKTQWN	NVENLNRFER	PMDYYEDLKM	
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	GLRAEGHPDS	LKDNSSCSVM	SEEPEGRSYS	TLTTVREIET	QT...ELLSP	
	<del>p</del>	<del>n</del>	<del>Eepe</del>	<del>e</del>	<del>l</del>	
	<u>P</u>		<u>EE</u>	<u>E</u>		
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	551					597
	ISRRVYV~	~~~~~	~~~~~	~~~~~	~~~~~	
	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~	
	DEAEARQDGY	GDRTLGYQYD	PEQLDLAENM	VSQNDGSFIS	KKEWYV~	
	GM.KFVSDEH	YDENEDDLVS	HV...DGSVI	SR...REWYV	~~~~~	
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	GSGRAEEEEED	QDEGIKQAMN	HFVQENGTLR	AKPTGNGIYI	NGRGHLV	
	g	A	E	DE	I	H
	<u>A</u>	<u>D</u>				<u>y</u>

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